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RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/601,561

DATE: 05/24/2001

TIME: 13:38:27

Input Set : A:\Nih332-1.app

Output Set: C:\CRF3\05242001\I601561.raw

ENTERED

3 <110> APPLICANT: Schneider, Thomas D.
 4 Hengen, Paul N.
 5 The Government of the United States of America
 6 as represented by The Secretary of the
 7 Department of Health and Human Services
 9 <120> TITLE OF INVENTION: Molecular Computing Elements: Gates and Flip-Flops
 11 <130> FILE REFERENCE: 015280-332100US
 13 <140> CURRENT APPLICATION NUMBER: US 09/601,561
 C--> 14 <141> CURRENT FILING DATE: 2001-05-14
 16 <150> PRIOR APPLICATION NUMBER: US 60/075,468
 17 <151> PRIOR FILING DATE: 1998-02-20
 19 <150> PRIOR APPLICATION NUMBER: WO PCT/US99/03469
 20 <151> PRIOR FILING DATE: 1999-02-17
 22 <160> NUMBER OF SEQ ID NOS: 19
 24 <170> SOFTWARE: PatentIn Ver. 2.1
 26 <210> SEQ ID NO: 1
 27 <211> LENGTH: 20
 28 <212> TYPE: DNA
 29 <213> ORGANISM: Artificial Sequence
 (31 <220> FEATURE:
 32 <223> OTHER INFORMATION: Description of Artificial Sequence:consensus
 33 sequence of early model of Factor for Inversion
 34 Stimulation (Fis) binding site
 36 <400> SEQUENCE: 1
 37 ttgstcaaaa ttgascaaa 20
 40 <210> SEQ ID NO: 2
 41 <211> LENGTH: 42
 42 <212> TYPE: DNA
 43 <213> ORGANISM: Artificial Sequence
 45 <220> FEATURE:
 46 <223> OTHER INFORMATION: Description of Artificial Sequence:paired Factor
 47 for Inversion Stimulation (Fis) binding sites with
 48 11 bp spacing; overlap 11
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 51 tattctttgc tcaaaatttg atcaaatttt gagcaaagaa ta 42
 54 <210> SEQ ID NO: 3
 55 <211> LENGTH: 38
 56 <212> TYPE: DNA
 57 <213> ORGANISM: Artificial Sequence
 59 <220> FEATURE:
 60 <223> OTHER INFORMATION: Description of Artificial Sequence:paired Factor
 61 for Inversion Stimulation (Fis) binding sites with
 62 7 bp spacing; overlap 7
 64 <400> SEQUENCE: 3
 65 aggcctttgc tcaaagtta aactttgagc aaaagcct 38
 68 <210> SEQ ID NO: 4
 69 <211> LENGTH: 15

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70 <212> TYPE: DNA
71 <213> ORGANISM: Artificial Sequence
73 <220> FEATURE:
74 <223> OTHER INFORMATION: Description of Artificial Sequence:sequence logo
75     for Factor for Inversion Stimulation (Fis) binding
76     site
78 <400> SEQUENCE: 4
79 gctcaaaaatt tgatc 15
82 <210> SEQ ID NO: 5
83 <211> LENGTH: 58
84 <212> TYPE: DNA
85 <213> ORGANISM: Artificial Sequence
87 <220> FEATURE:
88 <223> OTHER INFORMATION: Description of Artificial Sequence:Factor for
89     Inversion Stimulation (Fis) binding sites
90     separated by 23 bp; separated 23
92 <400> SEQUENCE: 5
93 ggaattcttt gctcaaaaatt tgatcaggat cctgatcaaa ttttgagcaa agaattcc 58
96 <210> SEQ ID NO: 6
97 <211> LENGTH: 21
98 <212> TYPE: DNA
99 <213> ORGANISM: Artificial Sequence
101 <220> FEATURE:
102 <223> OTHER INFORMATION: Description of Artificial Sequence:18.1 bit Fis
103     site
105 <400> SEQUENCE: 6
106 tttgctcaaa atttgatcaa a 21
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110 <211> LENGTH: 21
111 <212> TYPE: DNA
112 <213> ORGANISM: Artificial Sequence
114 <220> FEATURE:
115 <223> OTHER INFORMATION: Description of Artificial Sequence:18.1 bit Fis
116     site
118 <400> SEQUENCE: 7
119 tttgatcaaa ttttgagcaa a 21
122 <210> SEQ ID NO: 8
123 <211> LENGTH: 21
124 <212> TYPE: DNA
125 <213> ORGANISM: Artificial Sequence
127 <220> FEATURE:
128 <223> OTHER INFORMATION: Description of Artificial Sequence:12.7 bit Fis
129     site
131 <400> SEQUENCE: 8
132 tttgctcaaa gtttaaactt t 21
135 <210> SEQ ID NO: 9
136 <211> LENGTH: 21
137 <212> TYPE: DNA
138 <213> ORGANISM: Artificial Sequence

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140 <220> FEATURE:
141 <223> OTHER INFORMATION: Description of Artificial Sequence:12.7 bit Fis
142     site
144 <400> SEQUENCE: 9
145 aaagtttaaa ctttgagcaa a                                21
148 <210> SEQ ID NO: 10
149 <211> LENGTH: 21
150 <212> TYPE: DNA
151 <213> ORGANISM: Artificial Sequence
153 <220> FEATURE:
154 <223> OTHER INFORMATION: Description of Artificial Sequence:15.0 bit Fis
155     site
157 <400> SEQUENCE: 10
158 tttgctcaaa atttgatcag g                                21
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162 <211> LENGTH: 21
163 <212> TYPE: DNA
164 <213> ORGANISM: Artificial Sequence
166 <220> FEATURE:
167 <223> OTHER INFORMATION: Description of Artificial Sequence:15.0 bit Fis
168     site
170 <400> SEQUENCE: 11
171 cctgatcaaa ttttgagcaa a                                21
174 <210> SEQ ID NO: 12
175 <211> LENGTH: 46
176 <212> TYPE: DNA
177 <213> ORGANISM: Escherichia coli
179 <220> FEATURE:
180 <223> OTHER INFORMATION: origin of replication (oriC)
182 <400> SEQUENCE: 12
183 gttatacaca actcaaaaac tgaacaacag ttgttctttg gataac    46
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187 <211> LENGTH: 15
188 <212> TYPE: DNA
189 <213> ORGANISM: Artificial Sequence
191 <220> FEATURE:
192 <223> OTHER INFORMATION: Description of Artificial Sequence:Fis site
193     separated by 11 bases; 9.1 bit Fis site
195 <400> SEQUENCE: 13
196 gaacaacagt tgttc                                        15
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200 <211> LENGTH: 15
201 <212> TYPE: DNA
202 <213> ORGANISM: Artificial Sequence
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206     separated by 11 bases; 8.4 bit Fis site
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212 <210> SEQ ID NO: 15
213 <211> LENGTH: 113
214 <212> TYPE: DNA
215 <213> ORGANISM: Artificial Sequence
217 <220> FEATURE:
218 <223> OTHER INFORMATION: Description of Artificial Sequence:synthesized
219 single very long nucleic acid with hairpin loop
220 DNA
222 <400> SEQUENCE: 15
223 aacgggatcc actcaaaaac tgaacaacag ttgttcgaat tcctcgagcg atcggcgaag 60
224 ccgatcgctc gaggaattcg aacaactgtt gtccagtttt tgagtggatc ccg 113
227 <210> SEQ ID NO: 16
228 <211> LENGTH: 21
229 <212> TYPE: DNA
230 <213> ORGANISM: Artificial Sequence
232 <220> FEATURE:
233 <223> OTHER INFORMATION: Description of Artificial Sequence:8.4 bit Fis
234 site
236 <400> SEQUENCE: 16
237 tcactcaaa aactgaacaa c 21
240 <210> SEQ ID NO: 17
241 <211> LENGTH: 21
242 <212> TYPE: DNA
243 <213> ORGANISM: Artificial Sequence
245 <220> FEATURE:
246 <223> OTHER INFORMATION: Description of Artificial Sequence:10.0 bit Fis
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249 <400> SEQUENCE: 17
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253 <210> SEQ ID NO: 18
254 <211> LENGTH: 21
255 <212> TYPE: DNA
256 <213> ORGANISM: Artificial Sequence
258 <220> FEATURE:
259 <223> OTHER INFORMATION: Description of Artificial Sequence:10.0 bit Fis
260 site
262 <400> SEQUENCE: 18
263 ttcgaacaac tggtgttcag t 21
266 <210> SEQ ID NO: 19
267 <211> LENGTH: 21
268 <212> TYPE: DNA
269 <213> ORGANISM: Artificial Sequence
271 <220> FEATURE:
272 <223> OTHER INFORMATION: Description of Artificial Sequence:8.4 bit Fis
273 site
275 <400> SEQUENCE: 19
276 gttgttcagt ttttgagtg a 21

VERIFICATION SUMMARY

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L:14 M:271 C: Current Filing Date differs, Replaced Current Filing Date